



sequence listing.ST25.txt
SEQUENCE LISTING

<110> Cano, Carlos Antonio Durante
Nieto, Enrique Gerardo Guillen
Acosta, Anabel Álvarez
Munoz, Luis Emilio Carpio
Vazquez, Diogenes Quintana
Rodriguez, Carmen Elena Gomez
Rodriguez, Recardo de la Caridad Siva
Galvez, Consuelo Nazabal
Angulo, Maria de Jesus Leal
Dunn, Alejandro Miguel Martin

<120> Expression System of Heterologous Antigens as Fusion Proteins

<130> LEXSA P-13DIV2

<140> 09/612,925
<141> 2000-07-10

<150> US 08/930,917
<151> 1997-09-16

<150> PCT/CU97/00001
<151> 1997-01-17

<160> 31

<170> PatentIn version 3.2

<210> 1
<211> 1797
<212> DNA
<213> Neisseria meningitidis (group B)

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gtcaaagaag tttaaagttaa agtcggcgac aaaatctctg aagggtggtt gattgtcgtc 240
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gcgaaaagcc gtaaagtgga cgttatccaa ggcgacgggc aattcttaga tccgcaccac 720
ttggaagtgt cgctgactgc cggcgacgcg tacgaacagg cagcccctac cggcgagaaa 780
aaaatcgttg ccttcaaaaa ctgtatcatt gcagcaggca gccgcgtaac caaactgcct 840

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gaccgcgatt tggtaaaagt atggcaaaaa caaaacgaat accgttttga caacattatg   1080
gtcaacacca aaaccgttgc agttgagccg aaagaagacg gcgtttacgt tacctttgaa   1140
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gcgccaacg gcaaactcat cagcgcggaa aaagcaggcg ttgccgtaac cgatcgcggc   1260
ttcatcgaag tggacaaaca aatgcgtagc aatgtgccgc acatctacgc catcggcgac   1320
atcgtcggtc agccgatgtt ggcgcacaaa gccgttcacg aaggccacgt tgccgccgaa   1380
aactgcgccg gccacaaagc ctacttcgac gcacgcgtga ttccgggctg tgcctacact   1440
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atcaccaaag ccaacttccc gtgggcgggt tccggccgtg cgattgcaa cggttgcgac   1560
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gtcgggtccg acggtggcga tatgatcggc gaagtctgcc ttgccatcga aatgggctgc   1680
gacgcggcag acatcgga aaccatccac ccgcaccca ccttgggcga atccatcgggt   1740
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<210> 2
 <211> 47
 <212> PRT
 <213> Neisseria meningitidis (group B)

<400> 2

Met Val Asp Lys Arg Met Ala Leu Val Glu Leu Lys Val Pro Asp Ile
 1 5 10 15

Gly Gly His Glu Asn Val Asp Ile Ile Ala Val Glu Val Asn Val Gly
 20 25 30

Asp Thr Ile Ala Val Asp Asp Thr Leu Ile Thr Leu Asp Leu Glu
 35 40 45

<210> 3
 <211> 146
 <212> DNA
 <213> Neisseria meningitidis (group B)

<400> 3

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ttccatggta gataaaagaa tggcttagt tgaattgaaa gtgcccgaca ttggcggaca    60
cgaaaatgta gatattatcg cggttgaagt aaacgtgggc gacactattg ctgtggacga   120

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taccctgatt actttggatc tagaaa

146

<210> 4
 <211> 18
 <212> PRT
 <213> Neisseria meningitidis (group B)

<400> 4

Val Asn Val Gly Asp Thr Ile Ala Val Asp Asp Thr Leu Ile Thr Leu
 1 5 10 15

Asp Leu

<210> 5
 <211> 18
 <212> PRT
 <213> Neisseria meningitidis (group B)

<400> 5

Val Glu Val Gly Ser Lys Ile Tyr Val Asp Asp Gly Leu Ile Ser Leu
 1 5 10 15

Gln Val

<210> 6
 <211> 32
 <212> PRT
 <213> Neisseria meningitidis (group B)

<400> 6

Leu Val Glu Leu Lys Val Pro Asp Ile Gly Gly His Glu Asn Val Asp
 1 5 10 15

Ile Ile Ala Val Glu Val Asn Val Gly Asp Thr Ile Ala Val Asp Asp
 20 25 30

<210> 7
 <211> 32
 <212> PRT
 <213> Neisseria meningitidis (group B)

<400> 7

Leu Arg Glu Val Gln Val Pro Asp Arg Lys Leu His Lys Gly Val Gln
 1 5 10 15

Leu Leu Ala Gly Glu Leu Gly Ile Gly Glu Ala Leu Gln Val Asp Asp
 20 25 30

sequence listing.ST25.txt

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<210> 8
<211> 162
<212> PRT
<213> Neisseria meningitidis (group B)

<400> 8
Met Val Asp Lys Arg Met Ala Leu Val Glu Leu Lys Val Pro Asp Ile
1 5 10 15
Gly Gly His Glu Asn Val Asp Ile Ile Ala Val Glu Val Asn Val Gly
20 25 30
Asp Thr Ile Ala Val Asp Asp Thr Leu Ile Thr Leu Asp Leu Asp Ser
35 40 45
Arg Gly Ile Arg Ile Gly Pro Gly Arg Ala Ile Leu Ala Thr Ala Gly
50 55 60
Gly Gly Ala Arg Gln Ser Thr Pro Ile Gly Leu Gly Gly Ala Leu Tyr
65 70 75 80
Thr Thr Ala Gly Gly Gly Ala Arg Lys Ser Ile Thr Lys Gly Pro Gly
85 90 95
Arg Val Ile Tyr Ala Thr Ala Gly Gly Gly Ala Arg Lys Arg Ile His
100 105 110
Ile Gly Pro Gly Arg Ala Phe Tyr Thr Thr Ala Gly Gly Gly Ala Arg
115 120 125
Lys Arg Ile Thr Met Gly Pro Gly Arg Val Tyr Tyr Thr Thr Ala Gly
130 135 140
Gly Gly Ala Ser Ile Arg Ile Gln Arg Gly Pro Gly Arg Ala Phe Val
145 150 155 160

Thr Ile

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<210> 9
<211> 489
<212> DNA
<213> Neisseria meningitidis (group B)

<400> 9
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ctgattactt tggatctaga ctcgagaggc attcgtatcg gcccaggtcg cgcaatttta 180

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sequence listing.ST25.txt

gcaacagctg gcggtggcgc acgtcaatct acccctattg gtttaggtca ggctctgtat 240
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 gccaccgcgg gcgggcgggtgc ccgtaagcgt atccacattg gcccaggccg tgcattctat 360
 actacagcag gtggtggcgc acgtaaacgc atcactatgg gtcctgggtcg cgtctattac 420
 acgaccgctg gcgggcgggtgc tagcattcgc atccaacgcg gccctgggtcg tgcatttgtg 480
 accatatga 489

<210> 10
 <211> 47
 <212> PRT
 <213> Neisseria meningitidis (group B)

<400> 10

Met Leu Asp Lys Arg Met Ala Leu Val Glu Leu Lys Val Pro Asp Ile
 1 5 10 15

Gly Gly His Glu Asn Val Asp Ile Ile Ala Val Glu Val Asn Val Gly
 20 25 30

Asp Thr Ile Ala Val Asp Asp Thr Leu Ile Thr Leu Glu Thr Asp
 35 40 45

<210> 11
 <211> 29
 <212> DNA
 <213> Artificial

<220>
 <223> Primer 5' No. 1573

<400> 11
 ttccatggta gataaaagaa tggctttag 29

<210> 12
 <211> 29
 <212> DNA
 <213> Artificial

<220>
 <223> Primer 3' No. 1575

<400> 12
 tttctagatc caaagtaatc agggatcgc 29

<210> 13
 <211> 26
 <212> DNA
 <213> Artificial

<220>
 <223> Primer 3' No. 2192

sequence listing.ST25.txt

<400> 13
ggcggttctg ccgattaagg atccga 26

<210> 14
<211> 146
<212> DNA
<213> Artificial

<220>
<223> Derived fragment from the first 47 amino acids of the P64k antigen of N. meningitidis. The restriction sites NcoI (positions 3 to 8) and XbaI (positions 139 to 144) are introduced by PCR, which provokes changes in the nucleotide sequence of this

<400> 14
ttccatggta gataaaagaa tggctttagt tgaattgaaa gtgcccgaca ttggcggaca 60
cgaaaatgta gatattatcg cggttgaagt aaacgtgggc gacactattg ctgtggacga 120
taccctgatt actttggatc tagaaa 146

<210> 15
<211> 47
<212> PRT
<213> Artificial

<220>
<223> stabilizer fragment derived from the first 47 amino acids of the P64k antigen of N. meningitidis

<400> 15
Met Val Asp Lys Arg Met Ala Leu Val Glu Leu Lys Val Pro Asp Ile
1 5 10 15
Gly Gly His Glu Asn Val Asp Ile Ile Ala Val Glu Val Asn Val Gly
20 25 30
Asp Thr Ile Ala Val Asp Asp Thr Leu Ile Thr Leu Asp Leu Glu
35 40 45

<210> 16
<211> 16
<212> DNA
<213> Artificial

<220>
<223> oligonucleotide used to introduce restriction sites XbaI, EcoV, and BamHI in the 3' end of the stabilizer fragment of SEQ. ID. NO. 13

<400> 16
ctagatttga tatcag 16

<210> 17

sequence listing.ST25.txt

<211> 16
 <212> DNA
 <213> Artificial

<220>
 <223> Oligonucleotide used to introduce restriction sites XbaI, EcoV,
 and BamHI in the 3' end of the stabilizer fragment of SEQ. ID.
 NO. 13

<400> 17
 gatcctgata tcaaat

16

<210> 18
 <211> 15
 <212> PRT
 <213> Human immunodeficiency virus type 1

<400> 18

Ser Arg Gly Ile Arg Ile Gly Pro Gly Arg Ala Ile Leu Ala Thr
 1 5 10 15

<210> 19
 <211> 15
 <212> PRT
 <213> Human immunodeficiency virus type 1

<400> 19

Arg Gln Ser Thr Pro Ile Gly Leu Gly Gln Ala Leu Tyr Thr Thr
 1 5 10 15

<210> 20
 <211> 15
 <212> PRT
 <213> Human immunodeficiency virus type 1

<400> 20

Arg Lys Ser Ile Thr Lys Gly Pro Gly Arg Val Ile Tyr Ala Thr
 1 5 10 15

<210> 21
 <211> 15
 <212> PRT
 <213> Human immunodeficiency virus type 1

<400> 21

Arg Lys Arg Ile His Ile Gly Pro Gly Arg Ala Phe Tyr Thr Thr
 1 5 10 15

<210> 22
 <211> 15
 <212> PRT
 <213> Human immunodeficiency virus type 1

<400> 22

sequence listing.ST25.txt

Arg Lys Arg Ile Thr Met Gly Pro Gly Arg Val Tyr Tyr Thr Thr
1 5 10 15

<210> 23
<211> 15
<212> PRT
<213> Neisseria meningitidis (group B)

<400> 23

Ser Ile Arg Ile Gln Arg Gly Pro Gly Arg Ala Phe Val Thr Ile
1 5 10 15

<210> 24
<211> 15
<212> PRT
<213> Human immunodeficiency virus type 1

<400> 24

Thr Ser Ile Thr Ile Gly Pro Gly Gln Val Phe Tyr Arg Thr Gly
1 5 10 15

<210> 25
<211> 15
<212> PRT
<213> Human immunodeficiency virus type 1

<400> 25

Arg Gln Arg Thr Ser Ile Gly Gln Gly Gln Ala Leu Tyr Thr Thr
1 5 10 15

<210> 26
<211> 5
<212> PRT
<213> Artificial

<220>
<223> Spacer peptide that divides the various V3 epitopes in the MEPS
TAB3, TAB4, TAB9, and TAB13

<400> 26

Ala Gly Gly Gly Ala
1 5

<210> 27
<211> 141
<212> PRT
<213> Artificial

<220>
<223> Multiepitopic polypeptides that includes several copies of the
central part of the variable region 3 of the gp120 protein of the
HIV-1

sequence listing.ST25.txt

<400> 27

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Met Ala Pro Thr Ser Ser Ser Thr Ala Gln Thr Gln Leu Gln Leu Glu
1          5          10          15

His Leu Leu Leu Asp Leu Gln Ile Phe Leu Ser Arg Gly Ile Arg Ile
          20          25          30

Gly Pro Gly Arg Ala Ile Leu Ala Thr Ala Gly Gly Gly Ala Arg Gln
          35          40          45

Ser Thr Pro Ile Gly Leu Gly Gly Ala Leu Tyr Thr Thr Ala Gly Gly
          50          55          60

Gly Ala Arg Lys Ser Ile Thr Lys Gly Pro Gly Arg Val Ile Tyr Ala
65          70          75          80

Thr Ala Gly Gly Gly Ala Arg Lys Arg Ile His Ile Gly Pro Gly Arg
          85          90          95

Ala Phe Tyr Thr Thr Ala Gly Gly Gly Ala Arg Lys Arg Ile Thr Met
          100          105          110

Gly Pro Gly Arg Val Tyr Tyr Thr Thr Ala Gly Gly Gly Ala Ser Ile
          115          120          125

Arg Ile Gln Arg Gly Pro Gly Arg Ala Phe Val Thr Ile
          130          135          140

```

<210> 28

<211> 162

<212> PRT

<213> Artificial

<220>

<223> Multiepitopic polypeptides that includes several copies of the central part of the variable region 3 of the gp120 protein of the HIV-1

<400> 28

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Met Val Asp Lys Arg Met Ala Leu Val Glu Leu Lys Val Pro Asp Ile
1          5          10          15

Gly Gly His Glu Asn Val Asp Ile Ile Ala Val Glu Val Asn Val Gly
          20          25          30

Asp Thr Ile Ala Val Asp Asp Thr Leu Ile Thr Leu Asp Leu Asp Ser
          35          40          45

Arg Gly Ile Arg Ile Gly Pro Gly Arg Ala Ile Leu Ala Thr Ala Gly
          50          55          60

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50

55

60

Gly Gly Ala Arg Gln Ser Thr Pro Ile Gly Leu Gly Gly Ala Leu Tyr
 65 70 75 80

Thr Thr Ala Gly Gly Gly Ala Arg Lys Ser Ile Thr Lys Gly Pro Gly
 85 90 95

Arg Val Ile Tyr Ala Thr Ala Gly Gly Gly Ala Arg Lys Arg Ile His
 100 105 110

Ile Gly Pro Gly Arg Ala Phe Tyr Thr Thr Ala Gly Gly Gly Ala Arg
 115 120 125

Lys Arg Ile Thr Met Gly Pro Gly Arg Val Tyr Tyr Thr Thr Ala Gly
 130 135 140

Gly Gly Ala Ser Ile Arg Ile Gln Arg Gly Pro Gly Arg Ala Phe Val
 145 150 155 160

Thr Ile

<210> 29
 <211> 202
 <212> PRT
 <213> Artificial

<220>
 <223> Multiepitopic polypeptides that include several copies of the
 central part of the variable region 3 of the gp120 protein of the
 HIV-1

<400> 29

Met Val Asp Lys Arg Met Ala Leu Val Glu Leu Lys Val Pro Asp Ile
 1 5 10 15

Gly Gly His Glu Asn Val Asp Ile Ile Ala Val Glu Val Asn Val Gly
 20 25 30

Asp Thr Ile Ala Val Asp Asp Thr Leu Ile Thr Leu Asp Leu Asp Ser
 35 40 45

Arg Gly Ile Arg Ile Gly Pro Gly Arg Ala Ile Leu Ala Thr Ala Gly
 50 55 60

Gly Gly Ala Arg Gln Ser Thr Pro Ile Gly Leu Gly Gln Ala Leu Tyr
 65 70 75 80

sequence listing.ST25.txt

Thr Thr Ala Gly Gly Gly Ala Arg Lys Ser Ile Thr Lys Gly Pro Gly
85 90 95

Arg Val Ile Tyr Ala Thr Ala Gly Gly Gly Ala Arg Lys Arg Ile His
100 105 110

Ile Gly Pro Gly Arg Ala Phe Tyr Thr Thr Ala Gly Gly Gly Ala Arg
115 120 125

Lys Arg Ile Thr Met Gly Pro Gly Arg Val Tyr Tyr Thr Thr Ala Gly
130 135 140

Gly Gly Ala Arg Gln Arg Thr Ser Ile Gly Gln Gly Gln Ala Leu Tyr
145 150 155 160

Thr Thr Ala Gly Gly Gly Ala Thr Ser Ile Thr Ile Gly Pro Gly Gln
165 170 175

Val Phe Tyr Arg Thr Gly Ala Gly Gly Gly Ala Ser Ile Arg Ile Gln
180 185 190

Arg Gly Pro Gly Arg Ala Phe Val Thr Ile
195 200

<210> 30
<211> 368
<212> DNA
<213> Artificial

<220>
<223> Synthetic fragment that codifies for MEP TAB9. Restriction sites
XbaI and BamHI are introduced.

<400> 30
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tgggtgcgcg aaaagtatca ccaaggggtcc aggccgcgtc atttacgcca ccgcgggcgg 180
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cgggatcc 368

<210> 31
<211> 5
<212> PRT
<213> Neisseria meningitidis (group B)
<400> 31

sequence listing.ST25.txt

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